# mda-load library v1.0.0 manual

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This library is used to load MDA files created by the saveData program that is part of EPICS.

# 1 Compiling

Compiling requires that you specify the **mda-load** library. This can be done directly by specifying the filename libmda-load.a and its path. If the library is in a standard search location for libraries, you can use the "-l mda-load" option with gcc or ld.

If you are using Solaris, you will also need to specify the Networking Services Library (nsl). This can be done with the "-1 nsl" option.

#### 2 Code

An MDA file structure has a defined structure that is mimicked by the data structures this library uses. There first is a header describing the scan(s), followed by the highest dimensional scan, then all the lower dimensional scans, and finally by an optional section of extra PVs.

#### 2.1 Functions

There are functions for loading the entire file or just parts of it. The function and structure definitions are in the include file mda-load.h.

In the following functions, fptr is a FILE pointer corresponding to an open MDA file, normally gotten from fopen() and removed by fclose(). It doesn't matter what position the pointer is at in the file when calling these functions.

```
struct mda_file *mda_load( FILE *fptr)
```

This function loads the entire MDA file at once, returning a pointer to an mda\_file structure.

```
void mda_unload( struct mda_file *mda)
```

This will free the memory occupied by the mda\_file to which mda points.

```
struct mda_header *mda_header_load( FILE *fptr)
```

This function loads the MDA file header, returning a pointer to a mda\_header structure.

```
void mda_header_unload( struct mda_header *header)
```

This will free the memory occupied by the mda\_header to which header points.

```
struct scan *mda_scan_load( FILE *fptr)
```

This function loads the entire scan structure (all dimensions) for an MDA file, returning a pointer to a mda\_scan structure. It is the same as mda\_subscan\_load(), where depth is zero and recursive is one.

struct mda\_scan \*mda\_subscan\_load( FILE \*fptr, int depth, int \*indices, int recursive)

This function loads only a part of an MDA file's scans, returning a pointer to the resulting mda\_scan structure. The depth refers to how many dimensions down into the scan the subscan is located (it has to be less than the number of dimensions in the file), and to the number of members in indices. The indices array contains the location of the subscan, starting with the highest dimensional index; if depth is zero, indices can be set to NULL. The recursive flag determines whether the lower dimensional scans will also be read (if any exist).

```
void mda_scan_unload( struct mda_scan *scan)
```

This will free the memory occupied by the mda\_scan to which scan points.

```
struct mda_extra *mda_extra_load( FILE *fptr)
```

This function attempts to load the extra PVs for an MDA file. If they exist, it returns a pointer to an mda\_extra structure, otherwise it returns NULL.

```
void mda_extra_unload( struct mda_extra *extra)
```

This will free the memory occupied by the mda\_extra to which extra points. It is safe to call this function if extra is set to NULL.

### 2.2 Example

The simplest way to use this library is to simply load the entire MDA file, using the template below. Then there are only two functions needed, mda\_load() and mda\_unload(), with the rest of the work coming from accessing the MDA data structure.

```
#include "mda-load.h"
...
FILE *fptr;
struct mda_file *mda;
...
if( (fptr = fopen(filename, "r")) == NULL)
        exit(1);
if( (mda = mda_load(fptr)) == NULL)
        exit(1);
fclose(fptr);
...
/* Access the mda structure here. */
...
mda_unload(mda);
...
```

### 3 Data

Once you have a pointer to the allocated mda\_file, you can access its members directly. The definition of the various data structures are in mda-load.h. Assuming that the data pointer is called mda, here's how you access the data.

#### 3.1 Header

```
(struct mda_header *) mda->header
  (float) header->version
  (long) header->scan_number
  (short) header->data_rank
  (long) header->dimensions[n] , [n] = [0] to [data_rank - 1]
  (short) header->regular
  (long) header->extra_pvs_offset
```

This section contains the global data values for the MDA file. version signifies the MDA format version, normally 1.3. scan\_number is the number assigned by saveData to the scan. data\_rank show the number of dimensions to the scan (for a 3-D scan, this is 3). The dimensions array (with data\_rank elements) contains the number of elements for each dimension of the scan; for a 3-D scan, dimensions[1] is the number of elements to the 2-D scans. regular signifies whether the dimensions of any of the scans were changed while the overall scan was running. extra\_pvs\_offset gives, for the section of extra PV's, the offset in bytes from the beginning of the file; if that section does not exist, this will be 0.

#### 3.2 Scans

```
(struct mda_scan *) mda->scan
     (short) scan->scan_rank
      (long)
              scan->requested_points
              scan->last_point
     (long *) scan->offsets
      (char *) scan->name
      (char *) scan->time
      (short) scan->number_positioners
      (short) scan->number_detectors
      (short) scan->number_triggers
      (struct mda_positioner *) scan->positioners[n] ,
                                [n] = [0] to [scan->number_positioners - 1]
            (short) positioners[n]->number
            (char *) positioners[n]->name
            (char *) positioners[n]->description
            (char *) positioners[n]->step_mode
            (char *) positioners[n]->unit
            (char *) positioners[n]->readback_name
            (char *) positioners[n]->readback_description
            (char *) positioners[n]->readback_unit
     (struct mda_detector * scan->detectors[n] ,
                                [n] = [0] to [scan->number_detectors - 1]
            (short) detectors[n]->number
            (char *) detectors[n]->name
            (char *) detectors[n]->description
            (char *) detectors[n]->unit
```

This section includes the scan data. It is also recursive in nature due to it being able to handle arbitrary dimensions.

#### 3.2.1 Structure

The overall structure for multidimensional files is dictated by scan\_rank and sub\_scans. As long as scan\_rank is greater than one, sub\_scans will not be NULL and will contain an array of the next lower dimensional scans (it will be NULL if mda\_subscan\_load() was used to retrieve the scan when the recursive parameter set to zero). For a multidimensional scan, this takes the form of a tree, since each sub-scan can also have its own sub-scans. For a higher dimensional scan, the values for the positioners and detectors apply to all scan with its sub\_scans.

Suppose a  $5\times8\times20$  scan, where you want to access the (3,7,x) 1-D scan, you would access it as  $mda->scan->sub\_scans[2]->sub\_scans[6]$ . However, if the scan was aborted,  $mda->scan->sub\_scans[2]$  or  $mda->scan->sub\_scans[2]->sub\_scans[6]$  might be NULL, depending on where the scan was aborted. Using last\_point can let you know the last "officially valid" sub-scan is  $sub\_scans[last\_point - 1]$ . The reason that I say "officially valid" is that another scan might exist at  $sub\_scans[last\_point]$ , as it was the scan in progress that was aborted; one can use this data, but should take care.

#### 3.2.2 Variables

As described before, scan\_rank is the dimensionality of this scan. requested\_points is how many points were wanted, while last\_point tells how many actually were finished. offsets is an array of requested\_points members, showing the distance from the beginning of the MDA file to the subscans; if the value is zero, then that scan does not exist. name is the name of the scanner in EPICS, while time is when this particular scan was started.

number\_positioners tells how many positioners are moved as part of this scan. The positioners array, holding number\_positioners members, has a description of each positioner and its readback. number is the internal number the scanRecord uses to identify this positioner, while name is what its called, and description describes it. step\_mode is how the scan determined what step to use: it can be linear, where the spacing between steps is equal; table, where the step positions are read from an array; or fly, where the step positions are read back during an on-the-fly scan. unit is the associated unit of the positioner. Similarly, for the readback, there is readback\_name, readback\_description, and readback\_unit.

The detector information is very similar to the positioners, as there is a detectors array with number\_detectors elements. For each detectors, there is also a number, name, description, and unit.

The trigger information is again similar to the positioners, with a triggers array with number\_triggers elements. Each trigger has a number and name associated to it, as well as a command, which is a value sent to name to trigger.

The positioner data values are held in an two dimensional array named positioners\_data. Since one can't allocate a two dimensional array directly, it's actually an array of pointers (corresponding to each

detector), pointing to arrays of more pointers (corresponding the the data). To access the 8th data point of the 12th detector, one would type (scan->positioners\_data[11])[7]; the parentheses are not optional.

The detector data values, in detectors\_data, is accessed similarly to the positioner data values.

The sub\_scans variable is used for accessing lower dimensional scans (if they exist). It's described in Sec. ??.

#### 3.3 Extra PV's

This section, which doesn't always exist (signified by extra being NULL), contains extra PV's recorded during the scan. number\_pvs is the number of PV's contained, with the PV's being held in an array pvs.

For each PV, there is the name string and description string. type lets you know what kind of data type it is, with the correspondence seen in Table 1. If type isn't DBR\_STRING, count gives the number of elements to the array and unit string gives the unit for the values. The values themselves are held in an array values.

Table 1	: ]	Extra	PV	data	type
---------	-----	-------	----	------	------

type name	type value	C type Description		
DBR_STRING	0	(char *)	zero-terminated string	
DBR_CTRL_CHAR	32	(char *)	byte array	
DBR_CTRL_SHORT	29	(short *)	short integer array	
DBR_CTRL_LONG	33	(long *)	long integer array	
DBR_CTRL_FLOAT	30	(float *)	floating-point array	
DBR_CTRL_DOUBLE	34	(double *)	double-precision floating-point array	

Accessing the values is done by setting pointer values to the correct type, according to type and Table 1. Suppose the third extra PV was of type DBR\_CTRL\_DOUBLE, and you wanted to access its fifth member, this could be done using ((double \*) pvs[2]->values)[4].

## 4 Basic Information Routines

It is possible to load only the basic information of an MDA file, saving load time and memory. This information includes everything in the header, as well as the detector, positioner, and trigger descriptions for each scan dimension. As this information is taken from the first scan of each dimension, there is an assumption that every other scan is correctly represented by the first scan of its dimensionality.

#### 4.1 Functions

```
struct mda_fileinfo *mda_info_load( FILE *fptr)
```

This function loads a fileinfo of the MDA file, returning a pointer to an mda\_fileinfo structure.

void mda\_info\_unload( struct mda\_fileinfo \*fileinfo)
This will free the memory occupied by the mda\_fileinfo to which fileinfo points.

#### 4.2 Structure

```
(struct mda_fileinfo *) fileinfo
      (float) fileinfo->version
              fileinfo->scan_number
      (long)
      (short) fileinfo->data_rank
              fileinfo->dimensions[n], [n] = [0] to [data_rank - 1]
      (short) fileinfo->regular
      (long)
              fileinfo->last_topdim_point
      (char *) fileinfo->time
      (struct mda_scaninfo *) fileinfo->scaninfos[n] , [n] = [0] to [data_rank - 1]
            (short) scaninfos[n]->scan_rank
                    scaninfos[n]->requested_points
            (long)
            (char *) scaninfos[n]->name
            (short) scaninfos[n]->number_positioners
            (short) scaninfos[n]->number_detectors
            (short) scaninfos[n]->number_triggers
            (struct mda_positioner *) scaninfos[n]->positioners[n] ,
                                      [n] = [0] to [scan->number_positioners - 1]
                  (short) positioners[n]->number
                  (char *) positioners[n]->name
                  (char *) positioners[n]->description
                  (char *) positioners[n]->step_mode
                  (char *) positioners[n]->unit
                  (char *) positioners[n]->readback_name
                  (char *) positioners[n]->readback_description
                  (char *) positioners[n]->readback_unit
            (struct mda_detector * scaninfos[n]->detectors[n] ,
                                      [n] = [0] to [scan->number_detectors - 1]
                  (short) detectors[n]->number
                  (char *) detectors[n]->name
                  (char *) detectors[n]->description
                  (char *) detectors[n]->unit
            (struct mda_trigger *) scaninfos[n]->triggers[n] ,
                                      [n] = [0] to [scan->numbers_triggers - 1]
                  (short) triggers[n]->number
                  (char *) triggers[n]->name
                  (float) triggers[n]->command
```

The structure of the mda\_fileinfo structure is nonrecursive, and is thus easier to access. The variables version, scan\_number, data\_rank, dimensions, and regular are the same as described in Section 3.1. The variable last\_topdim\_point is the number of completed scans in the highest dimensional scan, while time is the data and time of the start of the overall measurement. The information for each dimension's scans are in the scaninfos array. The variables contained in each entry is a subset of those found in the earlier describes scan, and all are described in in Section 3.2.2.